

SEQUENCE LISTING

<110> The Trustees of the University of Pennsylvania
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Wilson, James M.

<120> Methods of Generating Chimeric Adenoviruses and Uses For Such
Chimeric Adenoviruses

<130> UPN-P3067PCT

<150> US 10/465,302
<151> 2003-06-20

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<160> 18

<170> PatentIn version 3.2

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				510		
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acg tac ttc agc ctg ggc aac aag ttt agg aac ccc acg gtg gcc ccc	Thr Tyr Phe Ser Leu Gly Asn Lys Phe Arg Asn Pro Thr Val Ala Pro	545	550	555	560	18412
acc cac gac gtg acg acg gac cgg tcc cag cgg ctg acg ctg cgg ttc	Thr His Asp Val Thr Thr Asp Arg Ser Gln Arg Leu Thr Leu Arg Phe	565	570	575		18460
gtg ccc gtc gac cgc gag gac acc gcg tac tcg tac aaa gtg cgc ttc	Val Pro Val Asp Arg Glu Asp Thr Ala Tyr Ser Tyr Lys Val Arg Phe	580	585	590		18508
acg ctg gcc gtg ggc gac aac cgc gtg ctg gac atg gcc agc acg tac	Thr Leu Ala Val Gly Asp Asn Arg Val Leu Asp Met Ala Ser Thr Tyr	595	600	605		18556
ttt gac atc cgc ggc gtg ttg gac cgc ggt ccc agc ttc aaa ccc tac	Phe Asp Ile Arg Gly Val Leu Asp Arg Gly Pro Ser Phe Lys Pro Tyr	610	615	620		18604
tcc ggc acc gcc tac aac tcc ctg gcc ccc aag ggc gcc ccc aac ccg	Ser Gly Thr Ala Tyr Asn Ser Leu Ala Pro Lys Gly Ala Pro Asn Pro	625	630	635	640	18652
tca gaa tgg aag ggc tca gac aac aaa att agt gta aga ggt cag gct	Ser Glu Trp Lys Gly Ser Asp Asn Lys Ile Ser Val Arg Gly Gln Ala	645	650	655		18700
ccg ttt ttt agt aca tcc att aca aag gat ggt att caa gtg gcc act	Pro Phe Phe Ser Thr Ser Ile Thr Lys Asp Gly Ile Gln Val Ala Thr	660	665	670		18748
gat act tct agc gga gct gtg tat gct aaa aag gaa tat cag cct gaa	Asp Thr Ser Ser Gly Ala Val Tyr Ala Lys Lys Glu Tyr Gln Pro Glu	675	680	685		18796
cca caa gta ggg caa gaa caa tgg aac agc gaa gcc agt gat agt gat	Pro Gln Val Gly Gln Glu Gln Trp Asn Ser Glu Ala Ser Asp Ser Asp	690	695	700		18844
aaa gta gct ggt agg att cta aaa gac aca aca ccc atg ttc cct tgt	Lys Val Ala Gly Arg Ile Leu Lys Asp Thr Thr Pro Met Phe Pro Cys	705	710	715	720	18892
tac ggt tcc tac gcc aag ccc aca aat gaa cag ggg ggg caa ggc act	Tyr Gly Ser Tyr Ala Lys Pro Thr Asn Glu Gln Gly Gly Gln Gly Thr	725	730	735		18940
aat act gta gat ctg cag ttc ttt gcc tct tca tcg gct acc tct acg						18988

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Val Ile Pro Pro Thr Val Thr Pro Thr Thr Leu Trp Thr Thr Ala	1850 1855 1860
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Lys Val Arg Phe Thr Leu Ala Val Gly Asp Asn Arg Val Leu Asp Met
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Ala Ser Thr Tyr Phe Asp Ile Arg Gly Val Leu Asp Arg Gly Pro Ser
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Phe Lys Pro Tyr Ser Gly Thr Ala Tyr Asn Ser Leu Ala Pro Lys Gly
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 Page 127

420

425

430

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 Leu Ala Phe Asn Ala Pro Leu Thr Leu Gln Ser Asp Thr Leu Asn Leu
 100 105 110
 Ala Cys Asn Ala Pro Leu Thr Val Gln Asp Asn Arg Leu Gly Ile Thr
 115 120 125
 Tyr Asn Ser Pro Leu Thr Leu Gln Asn Ser Glu Leu Ala Leu Ala Val
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 Thr Pro Pro Leu Asp Thr Ala Asn Asn Thr Leu Ala Leu Lys Thr Ala
 145 150 155 160
 Arg Pro Ile Ile Thr Asn Ser Asn Asn Glu Leu Thr Leu Ser Ala Asp
 165 170 175
 Ala Pro Leu Asn Thr Ser Thr Gly Thr Leu Arg Leu Gln Ser Ala Ala
 180 185 190
 Pro Leu Gly Leu Val Asp Gln Thr Leu Arg Val Leu Phe Ser Asn Pro
 195 200 205
 Leu Tyr Leu Gln Asn Asn Phe Leu Ser Leu Ala Ile Glu Arg Pro Leu
 210 215 220
 Ala Leu Thr Thr Thr Gly Ser Met Ala Met Gln Ile Ser Gln Pro Leu
 225 230 235 240
 Lys Val Glu Asp Gly Ser Leu Ser Leu Ser Ile Glu Ser Pro Leu Asn
 245 250 255
 Leu Lys Asn Gly Asn Leu Thr Leu Gly Thr Gln Ser Pro Leu Thr Val
 260 265 270
 Thr Gly Asn Asn Leu Ser Leu Thr Thr Thr Ala Pro Leu Thr Val Gln
 275 280 285
 Asn Asn Ala Leu Ala Leu Ser Val Leu Leu Pro Leu Arg Leu Phe Asn

290

295

300

Asn Thr Ser Leu Gly Val Ala Phe Asn Pro Pro Ile Ser Ser Ala Asn
 305 310 315 320
 Asn Gly Leu Ser Leu Asp Ile Gly Asn Gly Leu Thr Leu Gln Tyr Asn
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 Arg Leu Val Val Asn Ile Gly Gly Gly Leu Gln Phe Asn Asn Gly Ala
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 Ile Thr Ala Ser Ile Asn Ala Ala Leu Pro Leu Gln Tyr Ser Asn Asn
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 Gln Leu Ser Leu Asn Ile Gly Gly Gly Leu Arg Tyr Asn Gly Thr Tyr
 370 375 380
 Lys Asn Leu Ala Val Lys Thr Asp Ser Phe Arg Gly Leu Glu Ile Asp
 385 390 395 400
 Ser Asn Gln Phe Leu Val Pro Arg Leu Gly Ser Gly Leu Lys Phe Asp
 405 410 415
 Gln Tyr Gly Tyr Ile Ser Val Ile Pro Pro Thr Val Thr Pro Thr Thr
 420 425 430
 Leu Trp Thr Thr Ala Asp Pro Ser Pro Asn Ala Thr Phe Tyr Asp Ser
 435 440 445
 Leu Asp Ala Lys Val Trp Leu Ala Leu Val Lys Cys Asn Gly Met Val
 450 455 460
 Asn Gly Thr Ile Ala Ile Lys Ala Leu Lys Gly Thr Leu Leu Gln Pro
 465 470 475 480
 Thr Ala Ser Phe Ile Ser Phe Val Met Tyr Phe Tyr Ser Asn Gly Thr
 485 490 495
 Arg Arg Thr Asn Tyr Pro Thr Phe Glu Asn Glu Gly Ile Leu Ala Ser
 500 505 510
 Ser Ala Thr Trp Gly Tyr Arg Gln Gly Asn Ser Ala Asn Thr Asn Val
 515 520 525
 Thr Ser Ala Val Glu Phe Met Pro Ser Ser Thr Arg Tyr Pro Val Asn
 530 535 540
 Lys Gly Thr Glu Val Gln Asn Met Glu Leu Thr Tyr Thr Phe Leu Gln
 545 550 555 560
 Gly Asp Pro Thr Met Ala Ile Ser Phe Gln Ala Ile Tyr Asn His Ala
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Leu Glu Gly Tyr Ser Leu Lys Phe Thr Trp Arg Val Arg Asn Arg Glu
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Arg Phe Asp Ile Pro Cys Cys Ser Phe Ser Tyr Ile Thr Glu Glu
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